

## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant : Wei-Qiang Gao  
Appl. No. : Unknown  
Filed : Herewith  
For : HAIR CELL DISORDERS

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SEQUENCE SUBMISSION STATEMENT

Assistant Commissioner for Patents  
Washington, D.C. 20231

Dear Sir:

A copy of the Sequence Listing in computer readable form as required by 37 C.F.R. § 1.821(e) is submitted herewith.

As required by 37 C.F.R. § 1.821(f), the Sequence Listing information recorded in computer readable form on the enclosed disk is identical to the written Sequence Listing in the application filed herewith.

Respectfully submitted,

KNOBBE, MARTENS, OLSON & BEAR, LLP

Dated: 4 MAY 2001

By: 

James J. Mullen III, Ph.D.

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# SEQUENCE LISTING

<110> Genentech, Inc.  
Gao, Wei-Qiang

<120> HAIR CELL DISORDERS

<130> GENENT.035VPC

<160> 14

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 669

<212> PRT

<213> Homo sapiens

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Lys Glu Gly Arg Gly Lys Gly Lys Gly Lys Lys Lys Glu Arg Gly Ser
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Gly Lys Lys Pro Glu Ser Ala Ala Gly Ser Gln Ser Pro Ala Leu Pro
      65             70             75             80
Pro Arg Leu Lys Glu Met Lys Ser Gln Glu Ser Ala Ala Gly Ser Lys
      85             90             95
Leu Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser Ser Leu Arg Phe
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Lys Trp Phe Lys Asn Gly Asn Glu Leu Asn Arg Lys Asn Lys Pro Gln
      115            120            125
Asn Ile Lys Ile Gln Lys Lys Pro Gly Lys Ser Glu Leu Arg Ile Asn
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Lys Ala Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys Lys Val Ile Ser
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Lys Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr Ile Val Glu Ser
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Asn Glu Ile Ile Thr Gly Met Pro Ala Ser Thr Glu Gly Ala Tyr Val
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      210            215            220
Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn Gly Gly Glu Cys
      225            230            235            240
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<222> (2)...(2008)

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10

15

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20

25

30

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Glu Ser Arg Pro Arg Arg Ala Leu Arg Leu Arg Arg Asp Val Arg Ala

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Gln Arg Arg Gln Arg Gln Arg Glu Gly Gln Glu Glu Gly Ala Arg Leu

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70

75

80

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Ser Pro Ile Glu Arg Asp Glu Lys Pro Gly Ile Gly Cys Arg Phe Gln

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90

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Thr Ser Pro Ser Val \* Asn Gln Phe \* Ile Leu Leu Ser Gln Ile

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Gln Val Val Gln Glu Trp Glu \* Ile Glu Ser Lys Lys Gln Thr Thr

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Lys Tyr Gln Asp Thr Lys Lys Ala Arg Glu Val Arg Thr Ser His \*

130

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Gln Ser Ile Thr Gly \* Phe Trp Arg Val Tyr Val Gln Ser Asp Gln

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150

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165

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180

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Val Phe Arg Val Ser His * Asn Ile Ser Ile His Arg Arg Ser Lys	
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tac ttc ttc atc tac atc tac atc cac cac tgg gac aag cca tct tgt	673
Tyr Phe Phe Ile Tyr Ile Tyr Ile His His Trp Asp Lys Pro Ser Cys	
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aaa atg tgc gga gaa gga gaa aac ttt ctg tgt gaa tgg agg gga gtg	721
Lys Met Cys Gly Glu Gly Glu Asn Phe Leu Cys Glu Trp Arg Gly Val	
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Leu His Gly Glu Arg Pro Phe Lys Pro Leu Glu Ile Leu Val Gln Val	
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cca acc tgg att cac tgg agc aag atg tac tga gaa tgt gcc cat gaa	817
Pro Thr Trp Ile His Trp Ser Lys Met Tyr * Glu Cys Ala His Glu	
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Ser Pro Lys Pro Arg Lys Gly Gly Gly Ala Val Pro Glu Glu Ser Ala	
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gac cat aac cgg cat ctg cat cgc cct cct tgt ggt cgg cat cat gtg	913
Asp His Asn Arg His Leu His Arg Pro Pro Cys Gly Arg His His Val	
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Pro Ser Ser Ala Glu Pro Ser Val * Thr Lys Gln Tyr Asp Glu His	
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Gly Glu Ser Ile Arg Ile * Lys Arg His Leu Gln * Ala Tyr Cys	
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tga gag aga agc aga gac atc ctt ttc cac cag tca cta tac ttc cac	1153
* Glu Arg Ser Arg Asp Ile Leu Phe His Gln Ser Leu Tyr Phe His	
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Ser Pro Ser Leu His Tyr Cys His Pro Asp Ser * Pro Gln Leu Glu	
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Gln Arg Thr His * Lys His Pro Phe Arg Lys Pro Leu Cys Asn Arg	
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Lys Arg Thr Ser * Trp His Arg Arg Pro Ser * Met * Gln Leu	
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Pro Gln Ala Cys Gln Arg Asn Pro * Phe Leu Pro Arg Leu Ser Ser	
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Cys Arg Phe Pro His Ala Lys Leu Pro Gln Ile Ala Pro Phe Gly Asn	
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Leu Glu Asp Ser Gly Gly * Gly Val * Asn Asp Pro Arg Val Arg	
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Ala Ser Pro Arg Ala Cys * Glu Thr Arg Gln * Pro Ala Gly Gln	
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Gln His Lys Leu Pro Glu Gln * Leu Arg Glu * Asn Arg Arg *	
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Lys Ser Arg * Arg Tyr Ala Phe Pro Gly His Thr Glu Pro Pro Gly	

590 595 600

agc cag tct tga ggc aac acc tgc ctt ccg cct ggc tga cag cag gac 1969  
 Ser Gln Ser \* Gly Asn Thr Cys Leu Pro Pro Gly \* Gln Gln Asp  
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taa ccc agc agg ccg ctt ctc gac aca gga aga aat cca ggccaggctg 2018  
 \* Pro Ser Arg Pro Leu Leu Asp Thr Gly Arg Asn Pro  
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Ser	Lys	Leu	Val	Leu	Arg	Cys	Glu	Thr	Ser	Ser	Glu	Tyr	Ser	Ser	Leu
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Arg	Phe	Lys	Trp	Phe	Lys	Asn	Gly	Asn	Glu	Leu	Asn	Arg	Lys	Asn	Lys
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Ala	Ser	Phe	Tyr	Lys	His	Leu	Gly	Ile	Glu	Phe	Met	Glu	Ala	Glu	Glu
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 Thr Pro Ala Arg Met Ser Pro Val Asp Phe His Thr Pro Ser Ser Pro  
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 Lys Ser Pro Pro Ser Glu Met Ser Pro Pro Val Ser Ser Met Thr Val  
 485 490 495  
 Ser Met Pro Ser Met Ala Val Ser Pro Phe Met Glu Glu Glu Arg Pro  
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 His Pro Gln Gln Phe Ser Ser Phe His His Asn Pro Ala His Asp Ser  
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 Glu Ser Glu Thr Glu Asp Glu Arg Val Gly Glu Asp Thr Pro Phe Leu  
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 Arg Leu Ala Asp Ser Arg Thr Asn Pro Ala Gly Arg Phe Ser Thr Gln  
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Ser Glu Arg Lys Glu Gly Arg Gly Lys Gly Lys Gly Lys Lys Lys Glu
          35             40             45

cga ggc tcc ggc aag aag ccg gag tcc gcg gcg ggc agc cag agc cca      191
Arg Gly Ser Gly Lys Lys Pro Glu Ser Ala Ala Gly Ser Gln Ser Pro
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gcc ttg cct ccc caa ttg aaa gag atg aaa agc cag gaa tcg gct gca      239
Ala Leu Pro Pro Gln Leu Lys Glu Met Lys Ser Gln Glu Ser Ala Ala
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ggg tcc aaa cta gtc ctt ccg tgt gaa acc agt tct gaa tac tcc tct      287
Gly Ser Lys Leu Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser Ser
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ctc aga ttc aag tgg ttc aag aat ggg aat gaa ttg aat cga aaa aac      335
Leu Arg Phe Lys Trp Phe Lys Asn Gly Asn Glu Leu Asn Arg Lys Asn
          100            105            110

aaa cca caa aat atc aag ata caa aaa aag cca ggc aag tca gaa ctt      383
Lys Pro Gln Asn Ile Lys Ile Gln Lys Lys Pro Gly Lys Ser Glu Leu
          115            120            125

cgc att aac aaa gca tca ctg gct gat tct gga gag tat atg tgc aaa      431
Arg Ile Asn Lys Ala Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys Lys
          130            135            140

gtg atc agc aaa tta gga aat gac agt gcc tct gcc aat atc acc atc      479
Val Ile Ser Lys Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr Ile
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Val Glu Ser Asn Glu Ile Ile Thr Gly Met Pro Ala Ser Thr Glu Gly
          160            165            170            175

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Ala Tyr Val Ser Ser Glu Ser Pro Ile Arg Ile Ser Val Ser Thr Glu
  
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His	Leu	Val	Lys	Cys	Ala	Glu	Lys	Glu	Lys	Thr	Phe	Cys	Val	Asn	Gly					
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ggg	gag	tgc	ttc	atg	gtg	aaa	gac	ctt	tca	aac	ccc	tcg	aga	tac	ttg	719				
Gly	Glu	Cys	Phe	Met	Val	Lys	Asp	Leu	Ser	Asn	Pro	Ser	Arg	Tyr	Leu					
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Cys	Lys	Cys	Pro	Asn	Glu	Phe	Thr	Gly	Asp	Arg	Cys	Gln	Asn	Tyr	Val					
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Glu	Leu	Tyr	Gln	Lys	Arg	Val	Leu	Thr	Ile	Thr	Gly	Ile	Cys	Ile	Ala					
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Leu	Leu	Val	Val	Gly	Ile	Met	Cys	Val	Val	Ala	Tyr	Cys	Lys	Thr	Lys					
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Gly Pro Arg Glu Cys Asn Ser Phe Leu Arg His Ala Arg Glu Thr Pro	
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Asp Ser Tyr Arg Asp Ser Pro His Ser Glu Arg Tyr Val Ser Ala Met	
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Thr Thr Pro Ala Arg Met Ser Pro Val Asp Phe His Thr Pro Ser Ser	
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Pro Lys Ser Pro Pro Ser Glu Met Ser Pro Pro Val Ser Ser Met Thr	
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Val Ser Met Pro Ser Met Ala Val Ser Pro Phe Met Glu Glu Glu Arg	
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His His Pro Gln Gln Phe Ser Ser Phe His His Asn Pro Ala His Asp	
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Ser Asn Ser Leu Pro Ala Ser Pro Leu Arg Ile Val Glu Asp Glu Glu	
545 550 555	
tat gaa acg acc caa gag tac gag cca gcc caa gag cct gtt aag aaa	1727
Tyr Glu Thr Thr Gln Glu Tyr Glu Pro Ala Gln Glu Pro Val Lys Lys	
560 565 570 575	
ctc gcc aat agc cgg cgg gcc aaa aga acc aag ccc aat ggc cac att	1775
Leu Ala Asn Ser Arg Arg Ala Lys Arg Thr Lys Pro Asn Gly His Ile	
580 585 590	
gct aac aga ttg gaa gtg gac agc aac aca agc tcc cag agc agt aac	1823
Ala Asn Arg Leu Glu Val Asp Ser Asn Thr Ser Ser Gln Ser Ser Asn	
595 600 605	
tca gag agt gaa aca gaa gat gaa aga gta ggt gaa gat acg cct ttc	1871
Ser Glu Ser Glu Thr Glu Asp Glu Arg Val Gly Glu Asp Thr Pro Phe	
610 615 620	

ctg ggc ata cag aac ccc ctg gca gcc agt ctt gag gca aca cct gcc 1919  
 Leu Gly Ile Gln Asn Pro Leu Ala Ala Ser Leu Glu Ala Thr Pro Ala  
 625 630 635  
  
 ttc cgc ctg gct gac agc agg act aac cca gca ggc cgc ttc tcg aca 1967  
 Phe Arg Leu Ala Asp Ser Arg Thr Asn Pro Ala Gly Arg Phe Ser Thr  
 640 645 650 655  
  
 cag gaa gaa atc cag gcc agg ctg tct agt gta att gct aac caa gac 2015  
 Gln Glu Glu Ile Gln Ala Arg Leu Ser Ser Val Ile Ala Asn Gln Asp  
 660 665 670  
  
 cct att gct gta taaaacctaa ataaacacat agattcacct gtaaaacttt 2067  
 Pro Ile Ala Val  
 675  
  
 attttatata ataaagtatt ccaccttaaa ttaaacaatt tattttatatt tagcagttct 2127  
 gcaaatagaa aacaggaaaa aaacttttat aaattaaata tatgtatgta aaaatgaaaa 2187  
 aaaaaaaaaa aa 2199

<210> 5  
 <211> 637  
 <212> PRT  
 <213> Homo sapiens

<400> 5  
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 Pro Ala Leu Pro Pro Gln Leu Lys Glu Met Lys Ser Gln Glu Ser Ala  
 35 40 45  
 Ala Gly Ser Lys Leu Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser  
 50 55 60  
 Ser Leu Arg Phe Lys Trp Phe Lys Asn Gly Asn Glu Leu Asn Arg Lys  
 65 70 75 80  
 Asn Lys Pro Gln Asn Ile Lys Ile Gln Lys Lys Pro Gly Lys Ser Glu  
 85 90 95  
 Leu Arg Ile Asn Lys Ala Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys  
 100 105 110  
 Lys Val Ile Ser Lys Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr  
 115 120 125  
 Ile Val Glu Ser Asn Glu Ile Ile Thr Gly Met Pro Ala Ser Thr Glu  
 130 135 140  
 Gly Ala Tyr Val Ser Ser Glu Ser Pro Ile Arg Ile Ser Val Ser Thr  
 145 150 155 160  
 Glu Gly Ala Asn Thr Ser Ser Ser Thr Ser Thr Ser Thr Thr Gly Thr  
 165 170 175  
 Ser His Leu Val Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn  
 180 185 190  
 Gly Gly Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr  
 195 200 205  
 Leu Cys Lys Cys Pro Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr  
 210 215 220  
 Val Met Ala Ser Phe Tyr Lys Ala Glu Glu Leu Tyr Gln Lys Arg Val

225		230		235		240
Leu Thr Ile Thr Gly	Ile Cys Ile Ala Leu Leu Val Val Gly	Ile Met				
	245	250	255			
Cys Val Val Ala Tyr Cys Lys Thr Lys Lys Gln Arg Lys Lys Leu His						
	260	265	270			
Asp Arg Leu Arg Gln Ser Leu Arg Ser Glu Arg Asn Asn Met Met Asn						
	275	280	285			
Ile Ala Asn Gly Pro His His Pro Asn Pro Pro Pro Glu Asn Val Gln						
	290	295	300			
Leu Val Asn Gln Tyr Val Ser Lys Asn Val Ile Ser Ser Glu His Ile						
	305	310	315			
Val Glu Arg Glu Ala Glu Thr Ser Phe Ser Thr Ser His Tyr Thr Ser						
	325	330	335			
Thr Ala His His Ser Thr Thr Val Thr Gln Thr Pro Ser His Ser Trp						
	340	345	350			
Ser Asn Gly His Thr Glu Ser Ile Leu Ser Glu Ser His Ser Val Ile						
	355	360	365			
Val Met Ser Ser Val Glu Asn Ser Arg His Ser Ser Pro Thr Gly Gly						
	370	375	380			
Pro Arg Gly Arg Leu Asn Gly Thr Gly Gly Pro Arg Glu Cys Asn Ser						
	385	390	395			
Phe Leu Arg His Ala Arg Glu Thr Pro Asp Ser Tyr Arg Asp Ser Pro						
	405	410	415			
His Ser Glu Arg Tyr Val Ser Ala Met Thr Thr Pro Ala Arg Met Ser						
	420	425	430			
Pro Val Asp Phe His Thr Pro Ser Ser Pro Lys Ser Pro Pro Ser Glu						
	435	440	445			
Met Ser Pro Pro Val Ser Ser Met Thr Val Ser Lys Pro Ser Met Ala						
	450	455	460			
Val Ser Pro Phe Met Glu Glu Glu Arg Pro Leu Leu Leu Val Thr Pro						
	465	470	475			
Pro Arg Leu Arg Glu Lys Lys Phe Asp His His Pro Gln Gln Phe Ser						
	485	490	495			
Ser Phe His His Asn Pro Ala His Asp Ser Asn Ser Leu Pro Ala Ser						
	500	505	510			
Pro Leu Arg Ile Val Glu Asp Glu Glu Tyr Glu Thr Thr Gln Glu Tyr						
	515	520	525			
Glu Pro Ala Gln Glu Pro Val Lys Lys Leu Ala Asn Ser Arg Arg Ala						
	530	535	540			
Lys Arg Thr Lys Pro Asn Gly His Ile Ala Asn Arg Leu Glu Val Asp						
	545	550	555			
Ser Asn Thr Ser Ser Gln Ser Ser Asn Ser Glu Ser Glu Thr Glu Asp						
	565	570	575			
Glu Arg Val Gly Glu Asp Thr Pro Phe Leu Gly Ile Gln Asn Pro Leu						
	580	585	590			
Ala Ala Ser Leu Glu Ala Thr Pro Ala Phe Arg Leu Ala Asp Ser Arg						
	595	600	605			
Thr Asn Pro Ala Gly Arg Phe Ser Thr Gln Glu Glu Ile Gln Ala Arg						
	610	615	620			
Leu Ser Ser Val Ile Ala Asn Gln Asp Pro Ile Ala Val						
	625	630	635			

<210> 6

<211> 2460

<212> DNA  
<213> Homo sapiens

<220>

<221> CDS

<222> (461) ... (2371)

<400> 6

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cgggctcgcg cggaggccag gagctgagcg gcggcggtg ccggacgatg ggagcgtgag 120
caggacggtg ataacctctc cccgatcggg ttgcgagggc gccgggcaga ggccaggacg 180
cgagccgcca gcggcgggac ccatcgacga cttcccgggg cgacaggagc agccccgaga 240
gccagggcga gcgcccgttc caggtggccg gaccgcccgc cgcgtccgcg ccgcgtctcc 300
tgcaggcaac gggagacgcc ccgcgcagc gcgagcgctt cagcgcggcc gctcgtcttc 360
cccatcgagg gacaaacttt tcccaaacc gatccgagcc cttggaccaa actcgcttgc 420
gccgagagcc gtccgcgtag agcgtccgt ctccggcgag atg tcc gag cgc aaa 475
                                Met Ser Glu Arg Lys
                                1         5
```

```
gaa ggc aga ggc aaa ggg aag ggc aag aag aag gag cga ggc tcc ggc 523
Glu Gly Arg Gly Lys Gly Lys Gly Lys Lys Lys Glu Arg Gly Ser Gly
                                10         15         20
```

```
aag aag ccg gag tcc gcg gcg ggc agc cag agc cca gcc ttg cct ccc 571
Lys Lys Pro Glu Ser Ala Ala Gly Ser Gln Ser Pro Ala Leu Pro Pro
                                25         30         35
```

```
caa ttg aaa gag atg aaa agc cag gaa tcg gct gca ggt tcc aaa cta 619
Gln Leu Lys Glu Met Lys Ser Gln Glu Ser Ala Ala Gly Ser Lys Leu
                                40         45         50
```

```
gtc ctt cgg tgt gaa acc agt tct gaa tac tcc tct ctc aga ttc aag 667
Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser Ser Leu Arg Phe Lys
                                55         60         65
```

```
tgg ttc aag aat ggg aat gaa ttg aat cga aaa aac aaa cca caa aat 715
Trp Phe Lys Asn Gly Asn Glu Leu Asn Arg Lys Asn Lys Pro Gln Asn
                                70         75         80         85
```

```
atc aag ata caa aaa aag cca ggg aag tca gaa ctt cgc att aac aaa 763
Ile Lys Ile Gln Lys Lys Pro Gly Lys Ser Glu Leu Arg Ile Asn Lys
                                90         95         100
```

```
gca tca ctg gct gat tct gga gag tat atg tgc aaa gtg atc agc aaa 811
Ala Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys Lys Val Ile Ser Lys
                                105        110        115
```

```
tta gga aat gac agt gcc tct gcc aat atc acc atc gtg gaa tca aac 859
Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr Ile Val Glu Ser Asn
                                120        125        130
```

```
gag atc atc act ggt atg cca gcc tca act gaa gga gca tat gtg tct 907
Glu Ile Ile Thr Gly Met Pro Ala Ser Thr Glu Gly Ala Tyr Val Ser
                                135        140        145
```

tca gag tct ccc att aga ata tca gta tcc aca gaa gga gca aat act	955
Ser Glu Ser Pro Ile Arg Ile Ser Val Ser Thr Glu Gly Ala Asn Thr	
150 155 160 165	
tct tca tct aca tct aca tcc acc act ggg aca agc cat ctt gta aaa	1003
Ser Ser Ser Thr Ser Thr Ser Thr Thr Gly Thr Ser His Leu Val Lys	
170 175 180	
tgt gcg gag aag gag aaa act ttc tgt gtg aat gga ggg gag tgc ttc	1051
Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn Gly Gly Glu Cys Phe	
185 190 195	
atg gtg aaa gac ctt tca aac ccc tcg aga tac ttg tgc aag tgc cca	1099
Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr Leu Cys Lys Cys Pro	
200 205 210	
aat gag ttt act ggt gat cgc tgc caa aac tac gta atg gcc agc ttc	1147
Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr Val Met Ala Ser Phe	
215 220 225	
tac aag gcg gag gag ctg tac cag aag aga gtg ctg acc ata acc ggc	1195
Tyr Lys Ala Glu Glu Leu Tyr Gln Lys Arg Val Leu Thr Ile Thr Gly	
230 235 240 245	
atc tgc atc gcc ctc ctt gtg gtc ggc atc atg tgt gtg gtg gcc tac	1243
Ile Cys Ile Ala Leu Leu Val Val Gly Ile Met Cys Val Val Ala Tyr	
250 255 260	
tgc aaa acc aag aaa cag cgg aaa aag ctg cat gac cgt ctt cgg cag	1291
Cys Lys Thr Lys Lys Gln Arg Lys Lys Leu His Asp Arg Leu Arg Gln	
265 270 275	
agc ctt cgg tct gaa cga aac aat atg atg aac att gcc aat ggg cct	1339
Ser Leu Arg Ser Glu Arg Asn Asn Met Met Asn Ile Ala Asn Gly Pro	
280 285 290	
cac cat cct aac cca ccc ccc gag aat gtc cag ctg gtg aat caa tac	1387
His His Pro Asn Pro Pro Pro Glu Asn Val Gln Leu Val Asn Gln Tyr	
295 300 305	
gta tct aaa aac gtc atc tcc agt gag cat att gtt gag aga gaa gca	1435
Val Ser Lys Asn Val Ile Ser Ser Glu His Ile Val Glu Arg Glu Ala	
310 315 320 325	
gag aca tcc ttt tcc acc agt cac tat act tcc aca gcc cat cac tcc	1483
Glu Thr Ser Phe Ser Thr Ser His Tyr Thr Ser Thr Ala His His Ser	
330 335 340	
act act gtc acc cag act cct agc cac agc tgg agc aac gga cac act	1531
Thr Thr Val Thr Gln Thr Pro Ser His Ser Trp Ser Asn Gly His Thr	
345 350 355	
gaa agc atc ctt tcc gaa agc cac tct gta atc gtg atg tca tcc gta	1579
Glu Ser Ile Leu Ser Glu Ser His Ser Val Ile Val Met Ser Ser Val	
360 365 370	

gaa aac agt agg cac agc agc cca act ggg ggc cca aga gga cgt ctt	1627
Glu Asn Ser Arg His Ser Ser Pro Thr Gly Gly Pro Arg Gly Arg Leu	
375 380 385	
aat ggc aca gga ggc cct cgt gaa tgt aac agc ttc ctc agg cat gcc	1675
Asn Gly Thr Gly Gly Pro Arg Glu Cys Asn Ser Phe Leu Arg His Ala	
390 395 400 405	
aga gaa acc cct gat tcc tac cga gac tct cct cat agt gaa agg tat	1723
Arg Glu Thr Pro Asp Ser Tyr Arg Asp Ser Pro His Ser Glu Arg Tyr	
410 415 420	
gtg tca gcc atg acc acc ccg gct cgt atg tca cct gta gat ttc cac	1771
Val Ser Ala Met Thr Thr Pro Ala Arg Met Ser Pro Val Asp Phe His	
425 430 435	
acg cca agc tcc ccc aaa tcg ccc cct tcg gaa atg tct cca ccc gtg	1819
Thr Pro Ser Ser Pro Lys Ser Pro Pro Ser Glu Met Ser Pro Pro Val	
440 445 450	
tcc agc atg acg gtg tcc aag cct tcc atg gcg gtc agc ccc ttc atg	1867
Ser Ser Met Thr Val Ser Lys Pro Ser Met Ala Val Ser Pro Phe Met	
455 460 465	
gaa gaa gag aga cct cta ctt ctc gtg aca cca cca agg ctg cgg gag	1915
Glu Glu Glu Arg Pro Leu Leu Leu Val Thr Pro Pro Arg Leu Arg Glu	
470 475 480 485	
aag aag ttt gac cat cac cct cag cag ttc agc tcc ttc cac cac aac	1963
Lys Lys Phe Asp His His Pro Gln Gln Phe Ser Ser Phe His His Asn	
490 495 500	
ccc gcg cat gac agt aac agc ctc cct gct agc ccc ttg agg ata gtg	2011
Pro Ala His Asp Ser Asn Ser Leu Pro Ala Ser Pro Leu Arg Ile Val	
505 510 515	
gag gat gag gag tat gaa acg acc caa gag tac gag cca gcc caa gag	2059
Glu Asp Glu Glu Tyr Glu Thr Thr Gln Glu Tyr Glu Pro Ala Gln Glu	
520 525 530	
cct gtt aag aaa ctc gcc aat agc cgg cgg gcc aaa aga acc aag ccc	2107
Pro Val Lys Lys Leu Ala Asn Ser Arg Arg Ala Lys Arg Thr Lys Pro	
535 540 545	
aat ggc cac att gct aac aga ttg gaa gtg gac agc aac aca agc tcc	2155
Asn Gly His Ile Ala Asn Arg Leu Glu Val Asp Ser Asn Thr Ser Ser	
550 555 560 565	
cag agc agt aac tca gag agt gaa aca gaa gat gaa aga gta ggt gaa	2203
Gln Ser Ser Asn Ser Glu Ser Glu Thr Glu Asp Glu Arg Val Gly Glu	
570 575 580	
gat acg cct ttc ctg ggc ata cag aac ccc ctg gca gcc agt ctt gag	2251
Asp Thr Pro Phe Leu Gly Ile Gln Asn Pro Leu Ala Ala Ser Leu Glu	



585	590	595	
gca aca cct gcc ttc cgc ctg gct gac agc agg act aac cca gca ggc			2299
Ala Thr Pro Ala Phe Arg Leu Ala Asp Ser Arg Thr Asn Pro Ala Gly			
600	605	610	
cgc ttc tcg aca cag gaa gaa atc cag gcc agg ctg tct agt gta att			2347
Arg Phe Ser Thr Gln Glu Glu Ile Gln Ala Arg Leu Ser Ser Val Ile			
615	620	625	
gct aac caa gac cct att gct gta taaaacctaa ataaacacat agattcacct			2401
Ala Asn Gln Asp Pro Ile Ala Val			
630	635		
gtaaaacttt attttatata ataaagtatt ccaccttaaa ttaaacaatt tatttttatt			2460

<210> 7

<211> 241

<212> PRT

<213> Homo sapiens

<400> 7

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Glu Arg Gly Ser Gly Lys Lys Pro Glu Ser Ala Ala Gly Ser Gln Ser			
	20	25	30
Pro Ala Leu Pro Pro Gln Leu Lys Glu Met Lys Ser Gln Glu Ser Ala			
	35	40	45
Ala Gly Ser Lys Leu Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser			
	50	55	60
Ser Leu Arg Phe Lys Trp Phe Lys Asn Gly Asn Glu Leu Asn Arg Lys			
65	70	75	80
Asn Lys Pro Gln Asn Ile Lys Ile Gln Lys Lys Pro Gly Lys Ser Glu			
	85	90	95
Leu Arg Ile Asn Lys Ala Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys			
	100	105	110
Lys Val Ile Ser Lys Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr			
	115	120	125
Ile Val Glu Ser Asn Glu Ile Ile Thr Gly Met Pro Ala Ser Thr Glu			
	130	135	140
Gly Ala Tyr Val Ser Ser Glu Ser Pro Ile Arg Ile Ser Val Ser Thr			
145	150	155	160
Glu Gly Ala Asn Thr Ser Ser Ser Thr Ser Thr Ser Thr Thr Gly Thr			
	165	170	175
Ser His Leu Val Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn			
	180	185	190
Gly Gly Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr			
	195	200	205
Leu Cys Lys Cys Pro Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr			
	210	215	220
Val Met Ala Ser Phe Tyr Ser Thr Ser Thr Pro Phe Leu Ser Leu Pro			
225	230	235	240
Glu			

<210> 8  
 <211> 1715  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> CDS  
 <222> (493)...(1215)

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cggcggcggc tgccggacga tgggagcgtg agcaggacgg tgataacctc tccccgatcg 180
ggttgcgagg gcgccgggca gaggccagga cgcgagccgc cagcggcggg acccatcgac 240
gacttcccgg ggcgacagga gcagccccga gagccagggc gagcgcccg tccaggtggc 300
cggaccgccc gccgcgtccg cgccgcgctc cctgcaggca acgggagacg cccccgcgca 360
gcgcgagcgc ctcagcgcg cgctcgctc tccccatcga gggacaaaact tttcccaaac 420
ccgatccgag cccttgacc aaactcgctt gcgccgagag ccgtccgct agagcgctcc 480
gtctccggcg ag atg tcc gag cgc aaa gaa ggc aga ggc aaa ggg aag ggc 531
          Met Ser Glu Arg Lys Glu Gly Arg Gly Lys Gly Lys Gly
                1                5                10

aag aag aag gag cga ggc tcc ggc aag aag ccg gag tcc gcg gcg ggc 579
Lys Lys Lys Glu Arg Gly Ser Gly Lys Lys Pro Glu Ser Ala Ala Gly
      15                20                25

agc cag agc cca gcc ttg cct ccc caa ttg aaa gag atg aaa agc cag 627
Ser Gln Ser Pro Ala Leu Pro Pro Gln Leu Lys Glu Met Lys Ser Gln
      30                35                40                45

gaa tcg gct gca ggt tcc aaa cta gtc ctt cgg tgt gaa acc agt tct 675
Glu Ser Ala Ala Gly Ser Lys Leu Val Leu Arg Cys Glu Thr Ser Ser
      50                55                60

gaa tac tcc tct ctc aga ttc aag tgg ttc aag aat ggg aat gaa ttg 723
Glu Tyr Ser Ser Leu Arg Phe Lys Trp Phe Lys Asn Gly Asn Glu Leu
      65                70                75

aat cga aaa aac aaa cca caa aat atc aag ata caa aaa aag cca ggg 771
Asn Arg Lys Asn Lys Pro Gln Asn Ile Lys Ile Gln Lys Lys Pro Gly
      80                85                90

aag tca gaa ctt cgc att aac aaa gca tca ctg gct gat tct gga gag 819
Lys Ser Glu Leu Arg Ile Asn Lys Ala Ser Leu Ala Asp Ser Gly Glu
      95                100                105

tat atg tgc aaa gtg atc agc aaa tta gga aat gac agt gcc tct gcc 867
Tyr Met Cys Lys Val Ile Ser Lys Leu Gly Asn Asp Ser Ala Ser Ala
     110                115                120                125

aat atc acc atc gtg gaa tca aac gag atc atc act ggt atg cca gcc 915
Asn Ile Thr Ile Val Glu Ser Asn Glu Ile Ile Thr Gly Met Pro Ala
     130                135                140

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tca act gaa gga gca tat gtg tct tca gag tct ccc att aga ata tca 963  
 Ser Thr Glu Gly Ala Tyr Val Ser Ser Glu Ser Pro Ile Arg Ile Ser  
 145 150 155

gta tcc aca gaa gga gca aat act tct tca tct aca tct aca tcc acc 1011  
 Val Ser Thr Glu Gly Ala Asn Thr Ser Ser Ser Thr Ser Thr Ser Thr  
 160 165 170

act ggg aca agc cat ctt gta aaa tgt gcg gag aag gag aaa act ttc 1059  
 Thr Gly Thr Ser His Leu Val Lys Cys Ala Glu Lys Glu Lys Thr Phe  
 175 180 185

tgt gtg aat gga ggg gag tgc ttc atg gtg aaa gac ctt tca aac ccc 1107  
 Cys Val Asn Gly Gly Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro  
 190 195 200 205

tcg aga tac ttg tgc aag tgc cca aat gag ttt act ggt gat cgc tgc 1155  
 Ser Arg Tyr Leu Cys Lys Cys Pro Asn Glu Phe Thr Gly Asp Arg Cys  
 210 215 220

caa aac tac gta atg gcc agc ttc tac agt acg tcc act ccc ttt ctg 1203  
 Gln Asn Tyr Val Met Ala Ser Phe Tyr Ser Thr Ser Thr Pro Phe Leu  
 225 230 235

tct ctg cct gaa taggagcatg ctcagttggg gctgctttct tgttgctgca 1255  
 Ser Leu Pro Glu  
 240

tctccctca gattccacct agagctagat gtgtcttacc agatctaata ttgactgcct 1315  
 ctgcctgtcg catgagaaca ttaacaaaag caattgtatt acttcctctg ttgcgcacta 1375  
 gttggctctg agatactaata aggtgtgtga ggctcggat gtttctggaa ttgatattga 1435  
 atgatgtgat acaaattgat agtcaatatac aagcagtga atatagataat aaaggcattt 1495  
 caaagtctca cttttattga taaaataaaaa atcattctac tgaacagtcc atcttcttta 1555  
 tacaatgacc acatcctgaa aagggtgttg ctaagctgta accgatatgc acttgaaatg 1615  
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<210> 9  
 <211> 420  
 <212> PRT  
 <213> Homo sapiens

<400> 9  
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 Pro Ala Leu Pro Pro Gln Leu Lys Glu Met Lys Ser Gln Glu Ser Ala  
 35 40 45  
 Ala Gly Ser Lys Leu Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser  
 50 55 60  
 Ser Leu Arg Phe Lys Trp Phe Lys Asn Gly Asn Glu Leu Asn Arg Lys  
 65 70 75 80  
 Asn Lys Pro Gln Asn Ile Lys Ile Gln Lys Lys Pro Gly Lys Ser Glu  
 85 90 95

Leu Arg Ile Asn Lys Ala Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys  
 100 105 110  
 Lys Val Ile Ser Lys Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr  
 115 120 125  
 Ile Val Glu Ser Asn Glu Ile Ile Thr Gly Met Pro Ala Ser Thr Glu  
 130 135 140  
 Gly Ala Tyr Val Ser Ser Glu Ser Pro Ile Arg Ile Ser Val Ser Thr  
 145 150 155 160  
 Glu Gly Ala Asn Thr Ser Ser Ser Thr Ser Thr Ser Thr Thr Gly Thr  
 165 170 175  
 Ser His Leu Val Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn  
 180 185 190  
 Gly Gly Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr  
 195 200 205  
 Leu Cys Lys Cys Pro Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr  
 210 215 220  
 Val Met Ala Ser Phe Tyr Lys Ala Glu Glu Leu Tyr Gln Lys Arg Val  
 225 230 235 240  
 Leu Thr Ile Thr Gly Ile Cys Ile Ala Leu Leu Val Val Gly Ile Met  
 245 250 255  
 Cys Val Val Ala Tyr Cys Lys Thr Lys Lys Gln Arg Lys Lys Leu His  
 260 265 270  
 Asp Arg Leu Arg Gln Ser Leu Arg Ser Glu Arg Asn Asn Met Met Asn  
 275 280 285  
 Ile Ala Asn Gly Pro His His Pro Asn Pro Pro Pro Glu Asn Val Gln  
 290 295 300  
 Leu Val Asn Gln Tyr Val Ser Lys Asn Val Ile Ser Ser Glu His Ile  
 305 310 315 320  
 Val Glu Arg Glu Ala Glu Thr Ser Phe Ser Thr Ser His Tyr Thr Ser  
 325 330 335  
 Thr Ala His His Ser Thr Thr Val Thr Gln Thr Pro Ser His Ser Trp  
 340 345 350  
 Ser Asn Gly His Thr Glu Ser Ile Leu Ser Glu Ser His Ser Val Ile  
 355 360 365  
 Val Met Ser Ser Val Glu Asn Ser Arg His Ser Ser Pro Thr Gly Gly  
 370 375 380  
 Pro Arg Gly Arg Leu Asn Gly Thr Gly Gly Pro Arg Glu Cys Asn Ser  
 385 390 395 400  
 Phe Leu Arg His Ala Arg Glu Thr Pro Asp Ser Tyr Arg Asp Ser Pro  
 405 410 415  
 His Ser Glu Arg  
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<210> 10

<211> 2431

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (496)...(1755)

<400> 10

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gccggcgagg agttccccga aacttggttg aactccgggc tcgcgcggag gccaggagct 120  
 gagcggcggc ggctgccgga cgatgggagc gtgagcagga cggtgataac ctctccccga 180  
 tcgggttgcg agggcgccgg gcagaggcca ggacgcgagc cgccagcggc gggacccatc 240  
 gacgacttcc cggggcgaca ggagcagccc cgagagccag ggcgagcgcc cgttccaggt 300  
 ggccggaccg cccgcgcgct ccgcgcgcgc ctccctgcag gcaacgggag acgccccgc 360  
 gcagcgcgag cgcctcagcg cggccgctcg ctctcccat cgagggacaa acttttccca 420  
 aaccgatcc gagcccttgg accaaactcg cctgcgccga gagccgtccg cgtagagcgc 480  
 tccgtctccg gcgag atg tcc gag cgc aaa gaa ggc aga ggc aaa ggg aag 531  
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                   1                  5                  10

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Gly Glu Cys Phe Met	Val Lys Asp Leu Ser	Asn Pro Ser Arg Tyr Leu		
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Cys Lys Cys Pro Asn	Glu Phe Thr Gly Asp	Arg Cys Gln Asn Tyr Val		
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agaccctttt aagaagattt aaagaaaaac cactcggccc tgagtgcggc gaggaccctg	300
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Pro Tyr Arg Ser Leu Thr Arg Arg Arg Asp Ala Glu Arg Arg Tyr Thr	
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Ser Ser Ser Ala Asp Ser Glu Glu Gly Lys Ala Pro Gln Lys Ser Tyr	
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Ser Ser Ser Glu Thr Leu Lys Ala Tyr Asp Gln Asp Ala Arg Leu Ala	
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Tyr Gly Ser Arg Val Lys Asp Ile Val Pro Gln Glu Ala Glu Glu Phe	
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Cys Arg Thr Gly Ala Asn Phe Thr Leu Arg Glu Leu Gly Leu Glu Glu	
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Thr	Arg	Ser	Gly	Arg	Ser	Ser	Cys	Leu	Ser	Ser	Arg	Ala	Asn	Ser	Asn		
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Leu	Thr	Leu	Thr	Asp	Thr	Glu	His	Glu	Asn	Thr	Glu	Thr	Asp	His	Pro		
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Gly	Gly	Leu	Gln	Asn	His	Ala	Arg	Leu	Arg	Thr	Pro	Pro	Pro	Pro	Leu		
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Ser	His	Ala	His	Thr	Pro	Asn	Gln	His	His	Ala	Ala	Ser	Ile	Asn	Ser		
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ctg	aac	cgg	ggc	aac	ttc	acg	ccg	agg	agc	aac	ccc	agc	ccg	gcc	ccc	978	
Leu	Asn	Arg	Gly	Asn	Phe	Thr	Pro	Arg	Ser	Asn	Pro	Ser	Pro	Ala	Pro		
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Thr	Asp	His	Ser	Leu	Ser	Gly	Glu	Pro	Pro	Ala	Gly	Gly	Ala	Gln	Glu		
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Pro	Ala	His	Ala	Gln	Glu	Asn	Trp	Leu	Leu	Asn	Ser	Asn	Ile	Pro	Leu		
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Glu	Thr	Arg	Asn	Leu	Gly	Lys	Gln	Pro	Phe	Leu	Gly	Thr	Leu	Gln	Asp		
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Asn	Leu	Ile	Glu	Met	Asp	Ile	Leu	Gly	Ala	Ser	Arg	His	Asp	Gly	Ala		
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Tyr	Ser	Asp	Gly	His	Phe	Leu	Phe	Lys	Pro	Gly	Gly	Thr	Ser	Pro	Leu		
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Phe	Cys	Thr	Thr	Ser	Pro	Gly	Tyr	Pro	Leu	Thr	Ser	Ser	Thr	Val	Tyr		
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Glu Val Ala Ala Glu Arg Ser Ser Ser Pro Ser Thr Gln Leu Ser Ala
10 15 20 25

```

```

gac cca tct ctt gat ggg ctt ccg gca gca gaa gac atg cca gag ccc 629
Asp Pro Ser Leu Asp Gly Leu Pro Ala Ala Glu Asp Met Pro Glu Pro
30 35 40

```

```

cag act gaa gat ggg aga acc cct gga ctc gtg ggc ctg gcc gtg ccc 677
Gln Thr Glu Asp Gly Arg Thr Pro Gly Leu Val Gly Leu Ala Val Pro
45 50 55

```

```

tgc tgt gcg tgc cta gaa gct gag cgc ctg aga ggt tgc ctc aac tca 725
Cys Cys Ala Cys Leu Glu Ala Glu Arg Leu Arg Gly Cys Leu Asn Ser
60 65 70

```

```

gag aaa atc tgc att gtc ccc atc ctg gct tgc ctg gtc agc ctc tgc 773
Glu Lys Ile Cys Ile Val Pro Ile Leu Ala Cys Leu Val Ser Leu Cys
75 80 85

```

```

ctc tgc atc gcc ggc ctc aag tgg gta ttt gtg gac aag atc ttt gaa 821
Leu Cys Ile Ala Gly Leu Lys Trp Val Phe Val Asp Lys Ile Phe Glu

```

90	95	100	105	
tat gac tct cct act cac ctt gac cct ggg ggg tta ggc cag gac cct				869
Tyr Asp Ser Pro Thr His Leu Asp Pro Gly Gly Leu Gly Gln Asp Pro				
	110	115	120	
att att tct ctg gac gca act gct gcc tca gct gtg tgg gtg tcg tct				917
Ile Ile Ser Leu Asp Ala Thr Ala Ser Ala Val Trp Val Ser Ser				
	125	130	135	
gag gca tac act tca cct gtc tct agg gct caa tct gaa agt gag gtt				965
Glu Ala Tyr Thr Ser Pro Val Ser Arg Ala Gln Ser Glu Ser Glu Val				
	140	145	150	
caa gtt aca gtg caa ggt gac aag gct gtt gtc tcc ttt gaa cca tca				1013
Gln Val Thr Val Gln Gly Asp Lys Ala Val Val Ser Phe Glu Pro Ser				
	155	160	165	
gcg gca ccg aca ccg aag aat cgt att ttt gcc ttt tct ttc ttg ccg				1061
Ala Ala Pro Thr Pro Lys Asn Arg Ile Phe Ala Phe Ser Phe Leu Pro				
	170	175	180	185
tcc act gcg cca tcc ttc cct tca ccc acc cgg aac cct gag gtg aga				1109
Ser Thr Ala Pro Ser Phe Pro Ser Pro Thr Arg Asn Pro Glu Val Arg				
	190	195	200	
acg ccc aag tca gca act cag cca caa aca aca gaa act aat ctc caa				1157
Thr Pro Lys Ser Ala Thr Gln Pro Gln Thr Thr Glu Thr Asn Leu Gln				
	205	210	215	
act gct cct aaa ctt tct aca tct aca tcc acc act ggg aca agc cat				1205
Thr Ala Pro Lys Leu Ser Thr Ser Thr Ser Thr Thr Gly Thr Ser His				
	220	225	230	
ctt gta aaa tgt gcg gag aag gag aaa act ttc tgt gtg aat gga ggg				1253
Leu Val Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn Gly Gly				
	235	240	245	
gag tgc ttc atg gtg aaa gac ctt tca aac ccc tcg aga tac ttg tgc				1301
Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr Leu Cys				
	250	255	260	265
aag tgc cca aat gag ttt act ggt gat cgc tgc caa aac tac gta atg				1349
Lys Cys Pro Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr Val Met				
	270	275	280	
gcc agc ttc tac agt acg tcc act ccc ttt ctg tct ctg cct gaa				1394
Ala Ser Phe Tyr Ser Thr Ser Thr Pro Phe Leu Ser Leu Pro Glu				
	285	290	295	
taggagcatg ctcagttggt gctgctttct tggtgctgca tctccctca gattccacct				1454
agagctagat gtgtcttacc agatctaata ttgactgcct ctgcctgtcg catgagaaca				1514
ttaacaaaag caattgtatt acttcctctg ttccgcgacta gttggctctg agataactaat				1574
aggtgtgtga ggctccggat gtttctggaa ttgatattga atgatgtgat acaaattgat				1634
agtcaatatc aagcagtga atatatgataat aaaggcattt caaagtctca cttttattga				1694

taaaataaaa atcattctac tgaacagtcc atcttcttta tacaatgacc acatcctgaa 1754  
aagggtgttg ctaagctgta accgatatgc acttgaaatg atggtaagtt aattttgatt 1814  
cagaatgtgt tatttgtcac aaataaacat aataaaagga aaaaaaaac ccgaattc 1872

<210> 14

<211> 296

<212> PRT

<213> Homo sapiens

<400> 14

Met	Glu	Ile	Tyr	Ser	Pro	Asp	Met	Ser	Glu	Val	Ala	Ala	Glu	Arg	Ser	1	5	10	15
Ser	Ser	Pro	Ser	Thr	Gln	Leu	Ser	Ala	Asp	Pro	Ser	Leu	Asp	Gly	Leu	20	25	30	
Pro	Ala	Ala	Glu	Asp	Met	Pro	Glu	Pro	Gln	Thr	Glu	Asp	Gly	Arg	Thr	35	40	45	
Pro	Gly	Leu	Val	Gly	Leu	Ala	Val	Pro	Cys	Cys	Ala	Cys	Leu	Glu	Ala	50	55	60	
Glu	Arg	Leu	Arg	Gly	Cys	Leu	Asn	Ser	Glu	Lys	Ile	Cys	Ile	Val	Pro	65	70	75	80
Ile	Leu	Ala	Cys	Leu	Val	Ser	Leu	Cys	Leu	Cys	Ile	Ala	Gly	Leu	Lys	85	90	95	
Trp	Val	Phe	Val	Asp	Lys	Ile	Phe	Glu	Tyr	Asp	Ser	Pro	Thr	His	Leu	100	105	110	
Asp	Pro	Gly	Gly	Leu	Gly	Gln	Asp	Pro	Ile	Ile	Ser	Leu	Asp	Ala	Thr	115	120	125	
Ala	Ala	Ser	Ala	Val	Trp	Val	Ser	Ser	Glu	Ala	Tyr	Thr	Ser	Pro	Val	130	135	140	
Ser	Arg	Ala	Gln	Ser	Glu	Ser	Glu	Val	Gln	Val	Thr	Val	Gln	Gly	Asp	145	150	155	160
Lys	Ala	Val	Val	Ser	Phe	Glu	Pro	Ser	Ala	Ala	Pro	Thr	Pro	Lys	Asn	165	170	175	
Arg	Ile	Phe	Ala	Phe	Ser	Phe	Leu	Pro	Ser	Thr	Ala	Pro	Ser	Phe	Pro	180	185	190	
Ser	Pro	Thr	Arg	Asn	Pro	Glu	Val	Arg	Thr	Pro	Lys	Ser	Ala	Thr	Gln	195	200	205	
Pro	Gln	Thr	Thr	Glu	Thr	Asn	Leu	Gln	Thr	Ala	Pro	Lys	Leu	Ser	Thr	210	215	220	
Ser	Thr	Ser	Thr	Thr	Gly	Thr	Ser	His	Leu	Val	Lys	Cys	Ala	Glu	Lys	225	230	235	240
Glu	Lys	Thr	Phe	Cys	Val	Asn	Gly	Gly	Glu	Cys	Phe	Met	Val	Lys	Asp	245	250	255	
Leu	Ser	Asn	Pro	Ser	Arg	Tyr	Leu	Cys	Lys	Cys	Pro	Asn	Glu	Phe	Thr	260	265	270	
Gly	Asp	Arg	Cys	Gln	Asn	Tyr	Val	Met	Ala	Ser	Phe	Tyr	Ser	Thr	Ser	275	280	285	
Thr	Pro	Phe	Leu	Ser	Leu	Pro	Glu	290	295										